

1648  
11/14

BIOTECHNOLOGY  
SYSTEMS  
BRANCH

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/701,080B  
Source: 1609  
Date Processed by STIC: 9/27/2002

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TECH CENTER 1600/290

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
  - 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
- FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

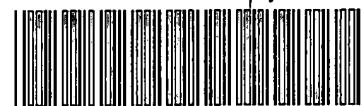
Revised 01/29/2002

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#14

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OCT 04 2002



TECH CENTER 1600/2900

1600

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/701,080B

DATE: 09/27/2002

TIME: 14:20:40

Input Set : A:\EP.txt

Output Set: N:\CRF4\09272002\I701080B.raw

3 <110> APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY  
 5 <120> TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN  
 P300 FOR USE IN  
 6 TRANSCRIPTIONAL REGULATION  
 8 <130> FILE REFERENCE: N73477C GCW  
 10 <140> CURRENT APPLICATION NUMBER: US 09/701080B  
 OK > 11 <141> CURRENT FILING DATE: 2001-02-27  
 13 <150> PRIOR APPLICATION NUMBER: GB 9811303.8  
 14 <151> PRIOR FILING DATE: 1998-05-26  
 16 <150> PRIOR APPLICATION NUMBER: GB 9900157.0  
 17 <151> PRIOR FILING DATE: 1999-01-05  
 19 <160> NUMBER OF SEQ ID NOS: 36  
 21 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply  
 Corrected Diskette Needed

ppr 1-2

## ERRORED SEQUENCES

24 <210> SEQ ID NO: 1  
 25 <211> LENGTH: 13 *12 shown (p.2)*  
 26 <212> TYPE: PRT  
 27 <213> ORGANISM: Artificial Sequence  
 29 <220> FEATURE:  
 30 <221> NAME/KEY: VARIANT  
 31 <222> LOCATION: (1)  
 32 <223> OTHER INFORMATION: Xaa represents Lys or Arg  
 34 <220> FEATURE:  
 35 <221> NAME/KEY: VARIANT  
 36 <222> LOCATION: (2)  
 37 <223> OTHER INFORMATION: Xaa represents Lys or Arg  
 39 <220> FEATURE:  
 40 <221> NAME/KEY: VARIANT  
 41 <222> LOCATION: (3)  
 42 <223> OTHER INFORMATION: Xaa represents any amino acid  
 44 <220> FEATURE:  
 45 <221> NAME/KEY: VARIANT  
 46 <222> LOCATION: (5)  
 47 <223> OTHER INFORMATION: Xaa represents any amino acid  
 49 <220> FEATURE:  
 50 <221> NAME/KEY: VARIANT  
 51 <222> LOCATION: (6)  
 52 <223> OTHER INFORMATION: Xaa represents any amino acid  
 54 <220> FEATURE:  
 55 <221> NAME/KEY: VARIANT  
 56 <222> LOCATION: (9)

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/701,080B

DATE: 09/27/2002  
TIME: 14:20:40

Input Set : A:\EP.txt  
Output Set: N:\CRF4\09272002\I701080B.raw

57 <223> OTHER INFORMATION: Xaa is Val or Ile  
59 <220> FEATURE:  
60 <221> NAME/KEY: VARIANT  
61 <222> LOCATION: (11)  
62 <223> OTHER INFORMATION: Xaa represents Lys or Arg  
64 <220> FEATURE:  
65 <221> NAME/KEY: VARIANT  
66 <222> LOCATION: (12)  
67 <223> OTHER INFORMATION: Xaa represents any amino acid  
69 <220> FEATURE:  
70 <223> OTHER INFORMATION: Description of Artificial Sequence: consensus sequence of  
transcriptional  
71 adaptor  
72 motif (TRAM)  
74 <400> SEQUENCE: 1 / / / / /  
W--> 75 Xaa Xaa Xaa Asn Xaa Xaa Cys Pro Xaa Cys Xaa Xaa  
E--> 76 1 5 10

Fug I

Use of n and/or Xaa has been detected in the Sequence Listing.  
Review the Sequence Listing to insure a corresponding  
explanation is presented in the <220> to <223> fields of  
each sequence using n or Xaa.

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/27/2002  
PATENT APPLICATION: US/09/701,080B TIME: 14:20:41

Input Set : A:\EP.txt  
Output Set: N:\CRF4\09272002\I701080B.raw

### Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 1,5,70

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/701,080B

DATE: 09/27/2002  
TIME: 14:20:41

Input Set : A:\EP.txt  
Output Set: N:\CRF4\09272002\I701080B.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:76 M:252 E: No. of Seq. differs, <211> LENGTH:Input:13 Found:12 SEQ:1  
L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:252 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0  
L:499 M:283 W: Missing Blank Line separator, <400> field identifier